Please amend the application as follows:

## In the specification:

Replace Table 1 on page 15 with the amended Table below.

Table 1. Nucleotide sequence of primers used to generate promoter fragments

Primer No.	Restriction site	Promoter sequence	Position	Sequence 5'-3'	
1582	HindIII (AAG CTT)	SEQ ID NO: 1	3440-3424	AAG CTT CTC GGC GCG CGG GCC C	Ğ
				(SEQ ID NO: 3)	
1583	Nhel (GCT AGC)	SEQ ID NO: 1	2341-2362	GCT AGC CAA GAG CTT CTG GAG C	:CG
	•.			(SEQ ID NO: 4)	* * * * * * * * * * * * * * * * * * *
1584	Nhel (GCT AGC)	SEQ ID NO: 1	720-741	GCT AGC TGT TAC ATG CAG AGC A	AT
-				(SEQ ID NO: 5)	
1585	HindIII (AAG CTT)	SEQ ID NO: 2	4439-4421	AAG CTT CCT ACG GCC CCC GCG	
				(SEQ ID NO: 6)	
1586	Nhel (GCT AGC)	SEQ ID NO: 2	3321-3340	GCT AGC GCG CAC TGC AAT GCC C	TC
			·	(SEQ ID NO: 7)	

Replace Table 2 on pages 20 and 21 with the amended Table below.

Table 2. Oligonucleotide primer used in the mutagenesis experiments

Primer		Primer Sequences
	<u>**</u>	
1949 P R1b Cre Fwd		CGCCGCCGT <b>TTG</b> GTCAGAGCCCCCT
	**	
		(SEQ ID NO: 8)
	<u> </u>	
1050		AGGGGGCTCTGAC <b>CAA</b> ACGGGCGGCG
1950 P R1b Cre Rev		
	:-	
		(SEQ ID NO: 9)
1951 P Rla GCI Fwd		CTCTCTTCCCCCCTAACTGCCTTCCC
	. ]	
		(SEQ ID NO: 10)
		GGGAAGGCAG <b>TTA</b> GGGGGGAAGAGAG
1952 P Rla GCI Rev		GGGAAGGCAGIIAGGGGGAAGAGAGA
	1 × 3	#:
		(SEQ ID NO: 11)
1953 P Rla GCII Fwd		GGCGGTCCAG <b>TTA</b> GGGGCTGGGATCC
	·	
		(SEQ ID NO: 12)
	( )	GGATCCCAGCCCC <b>TAA</b> CTGGACCGCC
1954 P Rla GCII Rev		
		(SEQ_ID_NO: 13)
2051 P Rla GCIII Fwd		CCTCTCCACCGCCCTAACCACCGCGCTGTG
TTO I TITU OUTII I WU	A.	332333333333333333333333333333333333333
		(SEQ ID NO: 14)
•		1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1
		CACAGCGCGGTGGTTAGGGCGGTGGAGAGG
2052 P Rla GCIII Rev		"CUCUAGO CAGTA CATTA CAGA CAGA CAGA CAGA CAGA CA
5035 & KTG GCTIT KGA		TORO ID NO. 151
		(SEQ ID NO: 15)

2053 P R1b GCIVs Fwd	•	CCCCAGCTCCCGCCCTAACCCCCCACCCC
	e egi	(SEQ ID NO: 16)
2054 P R1b GCIVs Rev		GGGGTGGGGGTTAGGGCGGGAGCTGGGG
		(600 TO NO 12)
	· ·	(SEQ ID NO: 17)
2055 P R1b GCV Fwd	· · · · · · · · · · · · · · · · · · ·	CGCTTCCCTCCCCTAACCCTTCCTGCC
		(SEQ ID NO: 18)
	·	(352 ID NO. 15)
2056 5 511 6611 5	•	GGCAGGAAGGGTTAGGGGAAGGG
2056 P R1b GCV Rev		
		(SEQ ID NO: 19)
2057 P R1b GCVI Fwd		CCCTCCCCTAACCTCCGACTGT
		(SEQ ID NO: 20)
2058 P R1b GCVI Rev		ACAGTCGGAGGTTAGGGGAGGGGAGGG
2000 I KID GOVI KEV		
		(SEQ ID NO: 21)
		·
2059 P R1b GCVII Fwd		CTCCGCCCACCCC <b>TAA</b> CTCCTGGCAC
		(SEQ ID NO: 22)
2060 P R1b GCVII Rev		GTGCCAGGAGTTAGGGGTGGGCGGAG
		(SEQ ID NO: 23)
2146 2 21	* *	
2146 P R1b GCIVd Fwd		CCCCAGCTCCCTAACTAACCCCCCACCCC
		(SEQ ID NO: 24)
		COCOMO COCOMO COMO COMO COMO COMO COMO
2147 P R1b GCIVd Rev		GGGGTGGGGGTTAGTTAGGGAGCTGGGG
		(SEQ ID NO: 25)

Replace the legend to Figure 6 running from page 25, line 24 through page 26, line 3 with the amended legend below.

Figure 6. Identification of nuclear factors binding to the Plb consensus CRE site using CREB/ATF super-shift antibodies.

Nuclear extracts (5µg) from ND7/23 cells were incubated with double-stranded <sup>32</sup>P-labeled oligonucleotides containing the Plb consensus CRE site (sense:5'-CGCCGCCCGTGACGTCAGAGCCCCCT-3' (SEQ ID NO: 26)). In lane 1, no antibody was added. In lane 2, a mouse monoclonal antibody (sc-270 Santa Cruz Biotechnology, Santa Cruz, CA) reactive with members of the ATF/CREB family such as ATF-1 p35, CREB-1 p43 and CREM-1 was pre-incubated at room temperature for 20 min before addition of <sup>32</sup>P-labeled probe. The specific complex between nuclear factors and the CRE is indicated by a star and the super-shifted complex is indicated by two stars.

Replace the present Sequence Listing with the revised one on 11 substitute sheets enclosed herewith.